



## SEQUENCE LISTING

<110> HONG, GUO FAN  
HUANG, WEI-HUA

<120> DNA POLYMERASE HAVING ABILITY TO REDUCE INNATE  
SELECTIVE DISCRIMINATION AGAINST FLUORESCENT  
DYE-LABELED DIDEOXYNUCLEOTIDES

<130> Lee109

<140> 09/512,019

<141> 2000-02-24

<150> 09/157,397

<151> 1998-09-12

<150> 08/642,684

<151> 1996-05-03

<160> 16

<170> PatentIn Ver. 3.2

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 agntgtttcc tgtgtgaaat tgttatccgn tcacaattcc acanaaaaata cgngncggnn 180  
 gnataaagt taaagcctgg ggtgnctaata gngtgngtta antcacatta attgngttgn 240  
 gntcaatgnc cgntttccag tcgggnaacc tgcgtgnca gntgnattaa tgggttcggcc 300  
 aacngncggg gngnggnggt ttgggtattg ggngntcttc cgnttcctcg ntcantgatt 360  
 cgttgngntc ggtcgttcgg ntgnggngng nggtatcaga tcantcaaag ggggtaatac 420  
 ggttatccac agaatacagg ggtaanggag gtaaggacat gtggggnaaa agggcagcaa 480  
 aagggcaggn accgtaaaaa ggccggttgg ttgggggttt tccatagggt ccgccccct 540  
 gggggggatc aaaaaaaatc cgnggccaaag tcaaggggtg gggggaccn ccagggnnta 600  
 taaagggtacc aggggttccc cctgggagtc cctccgtggg tctcctgtcc gccctgcccc 660  
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<210> 15  
 <211> 686  
 <212> DNA  
 <213> Unknown Organism

<220>  
 <223> Description of Unknown Organism: Template DNA  
 sequence

<220>  
 <221> modified\_base  
 <222> (561)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (656)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (678)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (680)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (682)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (685)  
 <223> a, c, g, t, other or unknown

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 tgagctaact cacattaatt gcgttgcgct cactgcccgc tttccagtcg ggaaacctgt 180  
 cgtgccagct gcattaatga atcggccaac gcgcggggag aggcgggttg cgtattgggc 240  
 gccagggtgg tttttctttt caccagttag acgggcaaca gctggattgc ccttcaccgc 300  
 ctggccctga gagagttaga gcaagcgggc cacgctgggt tgccccagca ggcgaaaatc 360  
 ctgtttgatg gtggttccga aatcggcaaa atcccttata aatcaaaaga ataggccgag 420  
 atagggttga gtgttgttcc agtttgggaa aagagtccac tattaagaa cgtggactcc 480  
 aacgtcaaag ggcgaaaaac cgtctatcag ggcgatgcca ctacgtgaac catcacccaa 540  
 atcaagtttt ttgggggtcga ngttgccgta aagcattaaa tcgggaacct aaaggagacc 600  
 ccgatttaga gcttgagggg gaaagccggc gaacgtgggc gagaaaaagg aggggnagaa 660  
 accgaaagga gcgcctnan gncgng 686

<210> 16  
 <211> 673  
 <212> DNA  
 <213> Unknown Organism

<220>  
 <223> Description of Unknown Organism: Template DNA  
 sequence

<220>  
 <221> modified\_base  
 <222> (9)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (27)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (41)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (468)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (491)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (564)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (591)  
 <223> a, c, g, t, other or unknown

<220>  
 <221> modified\_base  
 <222> (598)  
 <223> a, c, g, t, other or unknown

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 tagctgtttc ctgtgtgaaa ttgttatccg ctcaaatc cacacaacat acgagccgga 180  
 agcataaagt gtaaagcctg ggggtgcctaa tgagttagct aactcacatt aattgcgttg 240  
 cgctcactgc ccgctttcca gtcgggaaac ctgtcgtgcc agctgcatta atgaatcggc 300  
 caacgcgcgg ggagaggcgg ttgcggtatt gggcgctctt ccgcttcctc gctcactgac 360  
 tcgctgcgct cggctcgttcg gctgcggcgc gcggtatcag ctactcaaa ggcggtata 420  
 cggttatcca cagaatcagg ggataacgga ggtaaggaca tgtggggnaa aagggcagca 480  
 aaagggcagg naccgtaaaa aggcgggttg gttggggttt ttccataggg tccgcccccc 540  
 tgggggggat caaaaaaaaa cccngggccaa gtcaaggggt ggggggaccc nccagggntt 600  
 ataaagggtac caggggttcc ccctgggagt ccctccgtgg gtctcctgtc cgccctgccc 660  
 gttacccggt act 673